

Refine Search

Search Results -

Terms	Documents
L3 and (lung adj tumor)	256

Database:

US Pre-Grant Publication Full-Text Database
US Patents Full-Text Database
US OCR Full-Text Database
EPO Abstracts Database
JPO Abstracts Database
Derwent World Patents Index
IBM Technical Disclosure Bulletins

Search:

L4

[Refine Search]

Recall Text

Clear

Interrupt

Search History

DATE: Friday, June 09, 2006 [Printable Copy](#) [Create Case](#)

<u>Set Name</u>	<u>Query</u>	<u>Hit Count</u>	<u>Set Name</u>
result set			
<i>DB=PGPB,USPT,USOC,EPAB,JPAB,DWPI; PLUR=YES; OP=OR</i>			
<u>L4</u>	L3 and (lung adj tumor)	256	<u>L4</u>
<u>L3</u>	PRO831	261	<u>L3</u>
<i>DB=USPT; PLUR=YES; OP=OR</i>			
<u>L2</u>	PRO831	22	<u>L2</u>
<u>L1</u>	6951737.pn.	1	<u>L1</u>

END OF SEARCH HISTORY

SCORE Search Results Details for Application 10063534 and Search Result us-10-063-534- 30.rag.

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This page gives you Search Results detail for the Application 10063534 and Search Result us-10-063-534-30.rag.

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OM protein - protein search, using sw model

Run on: June 6, 2006, 21:30:04 ; Search time 200 Seconds
(without alignments)
166.884 Million cell updates/sec

Title: US-10-063-534-30
Perfect score: 386

Sequence: 1 MLLTLLLLLKGSCLEW.....PAMTCQAPRGEKVGDG 73

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 2589679 seqs, 457216429 residues

Total number of hits satisfying chosen parameters: 2589679

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : A_Geneseq_8:
1: geneseqp1980s:
2: geneseqp1990s:
3: geneseqp2000s:
4: geneseqp2001s:
5: geneseqp2002s:
6: geneseqp2003as:
7: geneseqp2003bs:
8: geneseqp2004s:
9: geneseqp2005s:
10: geneseqp2006s:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	386	100.0	73	3 AAY99346	Aay99346 Human PRO
2	386	100.0	73	4 AAB66095	Aab66095 Protein o
3	386	100.0	73	4 AAB87540	Aab87540 Human PRO
4	386	100.0	73	5 ABC95865	Abg95865 Human sec
5	386	100.0	73	6 ABU90890	Abu90890 Novel hum
6	386	100.0	73	6 ABC33949	Abo33949 Human sec
7	386	100.0	73	6 ABU71966	Abu71966 Novel hum
8	386	100.0	73	6 ABU71520	Abu71520 Human sec
9	386	100.0	73	6 ABU72301	Abu72301 Human PRO
10	386	100.0	73	6 ABU90974	Abu90974 Human PRO
11	386	100.0	73	6 ABO27295	Abo27295 Human sec
12	386	100.0	73	6 ABU92490	Abu92490 Human sec
13	386	100.0	73	6 ABU81160	Abu81160 Human sec
14	386	100.0	73	6 ABO53275	Abo53275 Novel hum
15	386	100.0	73	6 ABU98277	Abu98277 Novel hum
16	386	100.0	73	6 ABU89282	Abu89282 Novel hum
17	386	100.0	73	6 ABU82489	Abu82489 Novel hum
18	386	100.0	73	6 ABU96453	Abu96453 Human PRO

19	386	100.0	73	6	ABU72123	Abu72123 Human PRO
20	386	100.0	73	6	ADB17087	Adb17087 Human tra
21	386	100.0	73	6	ABO44253	Abo44253 Human sec
22	386	100.0	73	6	ABO33588	Abo33588 Novel hum
23	386	100.0	73	6	ADA19892	Ada19892 Novel hum
24	386	100.0	73	6	ADB17275	Adb17275 Human tra
25	386	100.0	73	6	ADA20064	Ada20064 Novel hum
26	386	100.0	73	6	ABO34181	Abo34181 Human sec
27	386	100.0	73	6	ADA00361	Ada00361 Human sec
28	386	100.0	73	7	ABO44441	Abo44441 Human sec
29	386	100.0	73	7	ABO33465	Abo33465 Novel hum
30	386	100.0	73	7	ADB85603	Adb85603 Novel hum
31	386	100.0	73	7	ADB68282	Adb68282 Human PRO
32	386	100.0	73	7	ADB68089	Adb68089 Human PRO
33	386	100.0	73	7	ADB90906	Adb90906 Novel hum
34	386	100.0	73	7	ADC06986	Adc06986 Human PRO
35	386	100.0	73	7	ADC17891	Adc17891 Human PRO
36	386	100.0	73	7	ADC17165	Adc17165 Mammalian
37	386	100.0	73	7	ADC14863	Adc14863 Novel hum
38	386	100.0	73	7	ADC52358	Adc52358 Novel hum
39	386	100.0	73	7	ADD70537	Add70537 Human sec
40	386	100.0	73	7	ADD39614	Add39614 Human sec
41	386	100.0	73	7	ADD70060	Add70060 Human sec
42	386	100.0	73	7	ADD36034	Add36034 Novel hum
43	386	100.0	73	7	ADD38181	Add38181 Human sec
44	386	100.0	73	7	ADD39137	Add39137 Human sec
45	386	100.0	73	7	ADD38660	Add38660 Human sec

ALIGNMENTS

RESULT 1
 AAY99346
 ID AAY99346 standard; protein; 73 AA.
 XX
 AC AAY99346;
 XX
 DT 08-AUG-2000 (first entry)
 XX
 DE Human PRO831 (UNQ471) amino acid sequence SEQ ID NO:22.
 XX
 KW Human; PRO polypeptide; membrane bound protein; receptor; diagnosis;
 KW transmembrane; secretion; immunoadhesion; pharmaceutical; screening.
 XX
 OS Homo sapiens.
 XX
 PN WO200012708-A2.
 XX
 PD 09-MAR-2000.
 XX
 PF 01-SEP-1999; 99WO-US020111.
 XX
 PR 01-SEP-1998; 98US-0098716P.
 PR 01-SEP-1998; 98US-0098749P.
 PR 01-SEP-1998; 98US-0098750P.
 PR 02-SEP-1998; 98US-0098803P.
 PR 02-SEP-1998; 98US-0098821P.
 PR 02-SEP-1998; 98US-0098843P.
 PR 09-SEP-1998; 98US-0099536P.
 PR 09-SEP-1998; 98US-0099536P.
 PR 09-SEP-1998; 98US-0099598P.
 PR 09-SEP-1998; 98US-0099602P.
 PR 09-SEP-1998; 98US-0099642P.
 PR 10-SEP-1998; 98US-0099741P.
 PR 10-SEP-1998; 98US-0099754P.
 PR 10-SEP-1998; 98US-0099763P.
 PR 10-SEP-1998; 98US-0099792P.
 PR 10-SEP-1998; 98US-0099808P.
 PR 10-SEP-1998; 98US-0099812P.
 PR 10-SEP-1998; 98US-0099815P.
 PR 10-SEP-1998; 98US-0099816P.
 PR 15-SEP-1998; 98US-0100385P.
 PR 15-SEP-1998; 98US-0100388P.
 PR 15-SEP-1998; 98US-0100390P.
 PR 16-SEP-1998; 98US-0100584P.
 PR 16-SEP-1998; 98US-0100627P.
 PR 16-SEP-1998; 98US-0100661P.
 PR 16-SEP-1998; 98US-0100662P.
 PR 16-SEP-1998; 98US-0100664P.
 PR 17-SEP-1998; 98US-0100683P.
 PR 17-SEP-1998; 98US-0100684P.
 PR 17-SEP-1998; 98US-0100710P.
 PR 17-SEP-1998; 98US-0100711P.
 PR 17-SEP-1998; 98US-0100919P.
 PR 17-SEP-1998; 98US-0100930P.
 PR 18-SEP-1998; 98US-0100848P.
 PR 18-SEP-1998; 98US-0100849P.
 PR 18-SEP-1998; 98US-0101014P.
 PR 18-SEP-1998; 98US-0101068P.
 PR 18-SEP-1998; 98US-0101071P.

SCORE Search Results Details for Application 10063534 and Search Result us-10-063-534- 30.olig.rag.

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OM protein - protein search, using sw model

Run on: June 6, 2006, 21:06:03 ; Search time 195 Seconds
(without alignments)
171.163 Million cell updates/sec

Title: US-10-063-534-30
Perfect score: 73

Sequence: 1 MLLLTLLLLLKGSCLEW.....PAMTCQAPRGEKVGDG 73

Scoring table: OLIGO
Gapop 60.0 , Gapext 60.0

Searched: 2589679 seqs, 457216429 residues

Word size : 6

Total number of hits satisfying chosen parameters: 27796

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

```
Database : A_Geneseq_8:*
1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
4: geneseqp2001s:*
5: geneseqp2002s:*
6: geneseqp2003as:*
7: geneseqp2003bs:*
8: geneseqp2004s:*
9: geneseqp2005s:*
10: geneseqp2006s:*
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8	73	100.0	73	ABU71520	Abu71520 Human sec
9	73	100.0	73	ABU72301	Abu72301 Human PRO
10	73	100.0	73	ABU90974	Abu90974 Human PRO
11	73	100.0	73	ABO27295	Abo27295 Human sec
12	73	100.0	73	ABU92490	Abu92490 Human sec
13	73	100.0	73	ABU81160	Abu81160 Human sec
14	73	100.0	73	ABO53275	Abo53275 Novel hum
15	73	100.0	73	ABU98277	Abu98277 Novel hum
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40	73	100.0	73	7	ADD39614	Add39614 Human sec
41	73	100.0	73	7	ADD70060	Add70060 Human sec
42	73	100.0	73	7	ADD36034	Add36034 Novel hum
43	73	100.0	73	7	ADD38181	Add38181 Human sec
44	73	100.0	73	7	ADD39137	Add39137 Human sec
45	73	100.0	73	7	ADD38660	Add38660 Human sec

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 PR 01-SEP-1998; 98US-0098750P.
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 PR 02-SEP-1998; 98US-0098821P.
 PR 02-SEP-1998; 98US-0098843P.
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 PR 10-SEP-1998; 98US-0099808P.
 PR 10-SEP-1998; 98US-0099812P.
 PR 10-SEP-1998; 98US-0099815P.
 PR 10-SEP-1998; 98US-0099816P.
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 PR 15-SEP-1998; 98US-0100390P.
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 PR 17-SEP-1998; 98US-0100711P.
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 PR 18-SEP-1998; 98US-0100849P.
 PR 18-SEP-1998; 98US-0101014P.
 PR 18-SEP-1998; 98US-0101068P.
 PR 18-SEP-1998; 98US-0101071P.